Report_a_defect/feature_request_for_caArray_in_Knowledge_Center

Feature Entry #1. Managing Vocabularies: If the users have to add their own vocabulary terms -and- they are to adhere to NCI or other ontology standards, then there should be a direct link to these vocabularies.

Feature Entry #2. LSD installation guide has configuration information for MySQL. It would be helpful to have reference in UPT installation to point to how to configure MySQL section in LSD installation guide.

Defect Entry #1 query anomalies. First, the user expects only two results, and there are 65. Second, Samples with no "id" attribute are returned, even you can see the query explicitly specifies an association to Sample objects where the id attribute is NOT NULL. There are results being returned for which the associated Sample has no "id" attribute. Third, the XML returned is fully populated, meaning the Experiment objects and their associations are returned to the user, which isn't something that is allowed in the CQL query specification, so I'm not sure why this is happening.

https://cabig-kc.nci.nih.gov/Molecular/forums/viewtopic.php?f=6&t=97&sid=ac8c97ac38e7785fda5a40658dcb4788

Defect Entry #2 "Manage Users" points to http://upt-dev.nci.nih.gov/upt/

 $\underline{https://cabig-kc.nci.nih.gov/Molecular/forums/viewtopic.php?f=6\&t=45\&sid=87bea40a114268c37dbdd7e414d0e87eacdefices.pdf.}$

Defect Entry #3 Uploading image files: caArray does not validate files in the SDRF Image File column. We have an open defect to address this shortcoming. Meanwhile, what you can do is to upload the image files as Supplemental Files.

 $\underline{https://cabig-kc.nci.nih.gov/Molecular/forums/viewtopic.php?f=6\&t=49\&sid=2e6073a51295b5a8bb5ca1f3eccbb188}$

Defect Entry #4 Certain file types can not be imported. The files will stay in ?Importing? state forever. The data files with these extension: .DAT, .TSV, .TIF haven?t been designated as either raw or derived, which makes the import step skip them.

Defect Entry #5 Wrong version on caGRID index service: We fixed domainModel.xml manually for 2.2.0. Project version field in the "domainModel.xml" was parsed incorrectly as an integer. As a remedy, a projectDescription of "Version 2.1 caArray Model" was added so that users can query this field to figure out the exact version

https://cabig-kc.nci.nih.gov/Molecular/forums/viewtopic.php?f=8&t=109

Defect Entry #6 Problem establishing socket connection from remote client. Internal Solution: https://cabig-kc.nci.nih.gov/Molecular/forums/viewtopic.php?f=8&t=91&sid=8dd14a09d59ca50221606904b7a8d64e it is part of 2.2.0

Defect Entry #7 geWorkbench-caArray slowness. Dan is investigating this; believed to be issue with graph cutting.